

re-run

RAW SEQUENCE LISTING

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Application Serial Number: 10/803,055
Source: FWO
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/803,055

DATE: 03/03/2005

TIME: 09:31:38

Input Set : N:\AMC\US10803055.raw

Output Set: N:\CRF4\03032005\J803055.raw

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1 <110> APPLICANT: Universite de Liege
2 <120> TITLE OF INVENTION: Cold-active beta galactosidase, the process for its
3   preparation and the use thereof
4 <130> FILE REFERENCE: Beta-gal
5 <140> CURRENT APPLICATION NUMBER: US/10/803,055
6 <141> CURRENT FILING DATE: 2004-03-18
7 <150> PRIOR APPLICATION NUMBER: US/09/501,136
8 <151> PRIOR FILING DATE: 2000-02-09
9 <160> NUMBER OF SEQ ID NOS: 2
10 <170> SOFTWARE: PatentIn Ver. 2.1
12 <210> SEQ ID NO: 1
13 <211> LENGTH: 3171
14 <212> TYPE: DNA
15 <213> ORGANISM: Pseudoalteromonas haloplanktis
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25   gaaggcgtaa actcggcttt tcatctttgg tgtaacgggc agtgggtggg gtattcacia 540
26   gatagccgct taccgagcga atttgattta agtgagcttt tagttgtcgg tactaaccgt 600
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31   tcaactgtcg agccgaaaa tcaaagcact aacaataaac gtgttgatga aaaagggggg 900
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36   catgctgtta gcaactgccg tatgattgaa gatattaagc tgatgaagca aaataacttt 1200
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47 tattgggctt acggcggcga ctttggatgat gaactaaacg accgccagtt ttgtataaac 1860
48 ggcttattgt tcccggatcg tacaccgcat cctagcctat ttgaagctaa atacagccag 1920
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71 <210> SEQ ID NO: 2

72 <211> LENGTH: 1039

73 <212> TYPE: PRT

74 <213> ORGANISM: Pseudoalteromonas haloplanktis

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77 <222> LOCATION: (460)

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79 <221> NAME/KEY: ACT_SITE

80 <222> LOCATION: (501)

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91 <221> NAME/KEY: SIMILAR

92 <222> LOCATION: (455) .. (460)

93 <400> SEQUENCE: 2

94 Met Thr Ser Leu Gln His Ile Ile Asn Arg Arg Asp Trp Glu Asn Pro

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95      1          5          10          15
96  Ile Thr Val Gln Val Asn Gln Val Lys Ala His Ser Pro Leu Asn Gly
97                20                25                30
98  Phe Lys Thr Ile Glu Asp Ala Arg Glu Asn Thr Gln Ser Gln Lys Lys
99                35                40                45
100  Ser Leu Asn Gly Gln Trp Asp Phe Lys Leu Phe Asp Lys Pro Glu Ala
101                50                55                60
102  Val Asp Glu Ser Leu Leu Tyr Glu Lys Ile Ser Lys Glu Leu Ser Gly
103                65                70                75                80
104  Asp Trp Gln Ser Ile Thr Val Pro Ser Asn Trp Gln Leu His Gly Phe
105                85                90                95
106  Asp Lys Pro Ile Tyr Cys Asn Val Lys Tyr Pro Phe Ala Val Asn Pro
107                100                105                110
108  Pro Phe Val Pro Ser Asp Asn Pro Thr Gly Cys Tyr Arg Thr Glu Phe
109                115                120                125
110  Thr Ile Thr Pro Glu Gln Leu Thr Gln Arg Asn His Ile Ile Phe Glu
111                130                135                140
112  Gly Val Asn Ser Ala Phe His Leu Trp Cys Asn Gly Gln Trp Val Gly
113                145                150                155                160
114  Tyr Ser Gln Asp Ser Arg Leu Pro Ser Glu Phe Asp Leu Ser Glu Leu
115                165                170                175
116  Leu Val Val Gly Thr Asn Arg Ile Ala Val Met Val Ile Arg Trp Ser
117                180                185                190
118  Asp Gly Ser Tyr Leu Glu Asp Gln Asp Met Trp Trp Leu Ser Gly Ile
119                195                200                205
120  Phe Arg Asp Val Asn Leu Leu Thr Lys Pro Gln Ser Gln Ile Arg Asp
121                210                215                220
122  Val Phe Ile Thr Pro Asp Leu Asp Ala Cys Tyr Arg Asp Ala Thr Leu
123                225                230                235                240
124  His Ile Lys Thr Ala Ile Asn Ala Pro Asn Asn Tyr Gln Val Ala Val
125                245                250                255
126  Gln Ile Phe Asp Gly Lys Thr Ser Leu Cys Glu Pro Lys Ile Gln Ser
127                260                265                270
128  Thr Asn Asn Lys Arg Val Asp Glu Lys Gly Gly Trp Ser Asp Val Val
129                275                280                285
130  Phe Gln Thr Ile Ala Ile Arg Ser Pro Lys Lys Trp Thr Ala Glu Thr
131                290                295                300
132  Pro Tyr Leu Tyr Arg Cys Val Val Ser Leu Leu Asp Glu Gln Gly Asn
133                305                310                315                320
134  Thr Val Asp Val Glu Ala Tyr Asn Ile Gly Phe Arg Lys Val Glu Met
135                325                330                335
136  Leu Asn Gly Gln Leu Cys Val Asn Gly Lys Pro Leu Leu Ile Arg Gly
137                340                345                350
138  Val Asn Arg His Glu His His Pro Glu Asn Gly His Ala Val Ser Thr
139                355                360                365
140  Ala Asp Met Ile Glu Asp Ile Lys Leu Met Lys Gln Asn Asn Phe Asn
141                370                375                380
142  Ala Val Arg Thr Ala His Tyr Pro Asn His Pro Leu Phe Tyr Glu Leu
143                385                390                395                400

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144   Cys Asp Glu Leu Gly Leu Tyr Val Val Asp Glu Ala Asn Ile Glu Thr
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146   His Gly Met Phe Pro Met Gly Arg Leu Ala Ser Asp Pro Leu Trp Ala
147           420                               425           430
148   Gly Ala Phe Met Ser Arg Tyr Thr Gln Met Val Glu Arg Asp Lys Asn
149           435                               440           445
150   His Ala Ser Ile Ile Ile Trp Ser Leu Gly Asn Glu Cys Gly His Gly
151           450                               455           460
152   Ala Asn His Asp Ala Met Tyr Gly Trp Ser Lys Ser Phe Asp Pro Ser
153           465                               470           475           480
154   Arg Pro Val Gln Tyr Glu Gly Gly Gly Ala Asn Thr Thr Ala Thr Asp
155           485                               490           495
156   Ile Ile Cys Pro Met Tyr Ser Arg Val Asp Thr Asp Ile Lys Asp Asp
157           500                               505           510
158   Ala Val Pro Lys Tyr Ser Ile Lys Lys Trp Leu Ser Leu Pro Gly Glu
159           515                               520           525
160   Thr Arg Pro Leu Ile Leu Cys Glu Tyr Ala His Ala Met Gly Asn Ser
161           530                               535           540
162   Leu Gly Ser Phe Asp Asp Tyr Trp Gln Ala Phe Arg Glu Tyr Pro Arg
163           545                               550           555           560
164   Leu Gln Gly Gly Phe Ile Trp Asp Trp Val Asp Gln Gly Leu Ser Lys
165           565                               570           575
166   Ile Asp Glu Asn Gly Lys His Tyr Trp Ala Tyr Gly Gly Asp Phe Gly
167           580                               585           590
168   Asp Glu Leu Asn Asp Arg Gln Phe Cys Ile Asn Gly Leu Leu Phe Pro
169           595                               600           605
170   Asp Arg Thr Pro His Pro Ser Leu Phe Glu Ala Lys Tyr Ser Gln Gln
171           610                               615           620
172   His Leu Gln Phe Thr Leu Arg Glu Gln Asn Gln Asn Gln Asn Gln Asn
173           625                               630           635           640
174   Gln Tyr Ser Ile Asp Val Phe Ser Asp Tyr Val Phe Arg His Thr Asp
175           645                               650           655
176   Asn Glu Lys Leu Val Trp Gln Leu Ile Gln Asn Gly Val Cys Val Glu
177           660                               665           670
178   Gln Gly Glu Met Ala Leu Asn Ile Ala Pro Gln Ser Thr His Thr Leu
179           675                               680           685
180   Thr Ile Lys Thr Lys Thr Ala Phe Glu His Gly Ala Gln Tyr Tyr Leu
181           690                               695           700
182   Asn Leu Asp Val Ala Leu Ile Asn Asp Ser His Phe Ala Asn Ala Asn
183           705                               710           715           720
184   His Val Met Asp Ser Glu Gln Phe Lys Leu Ile Asn Ser Asn Asn Leu
185           725                               730           735
186   Asn Ser Lys Ser Phe Ala Ser Ala Thr Glu Lys Ser Val Ile Ser Val
187           740                               745           750
188   Asn Glu Thr Asp Ser His Leu Ser Ile Glu Asn Asn Thr Phe Lys Leu
189           755                               760           765
190   Val Phe Asn Gln Gln Ser Gly Leu Ile Glu Gln Trp Leu Gln Asp Asp
191           770                               775           780
192   Thr Gln Val Ile Ser Ser Pro Leu Val Asp Asn Phe Tyr Arg Ala Pro

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198	Thr	Cys	Ser	Ser	Ile	Asn	Ala
199							
200	Thr	Cys	Val	Phe	Asn	Tyr	Glu
201							
202	Gln	Trp	Leu	Tyr	Thr	Leu	Asn
203							
204	Asp	Val	Asn	Leu	Asn	Asp	Thr
205							
206	Ser	Thr	Thr	Ile	Asn	Lys	Gln
207							
208	Leu	Gly	Pro	Phe	Glu	Asn	Tyr
209							
210	Gly	Tyr	Tyr	Ser	Leu	Ser	Leu
211							
212	Pro	Thr	Asp	Asn	Gly	Leu	Arg
213							
214	Asn	Leu	Ile	Val	Thr	Gly	Ala
215							
216	Gln	Asn	Met	Leu	Thr	Gln	Ala
217							
218	Asp	Cys	Ile	His	Val	His	Ile
219							
220	Asp	Asp	Ser	Trp	Ser	Pro	Ser
221							
222	Lys	Asn	Tyr	Asn	Tyr	Ser	Leu
223							

VERIFICATION SUMMARY

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